

Sayala

CRF Errors Corrected by the STIC Systems Branch

1761
11/17/98
CRF Processing Date: 11/17/98
Edited by: AJ
Verified by: AJ (STIC staff)

Serial Number: 08/631470A

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

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RAW SEQUENCE LISTING
PATENT APPLICATION US/08/631,470ADATE: 11/17/1998
TIME: 18:03:11

Sayala

Input Set: H631470A.RAW

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

Does Not Comply
Corrected Diskette Needed

1 <110> Brod, Staley A.
2 <120> Method of Treating Auto-Immune Diseases Using Type
3 One Interferons
4 <130> D5716CIP2
5 <140> US/08/631,470A
6 <141> 1996-04-12
7 <150> US 08/408,271
8 <151> 1995-03-24
9 <160> 6

ERRORED SEQUENCES FOLLOW

10 <210> 6
E--> 11 <211> 25
12 <212> DNA
13 <213> artificial sequence
14 <220>
15 <222> 687..663
16 <223> Oligonucleotide probe, designated by the relative
17 position in the human mRNA, used to determine the
18 specificity of the amplified GAPDH mRNA.
19 <400> 6
20 ggctctccag aacatcatcc ctgcc 25
E--> 21 seq 3

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VERIFICATION SUMMARY
PATENT APPLICATION US/08/631,470A

DATE: 11/17/1998
TIME: 18:03:11

Input Set: H631470A.RAW

Line	? Error/Warning	Original Text
11	E Input 25, Calc# Bases 28 differ	<211> 25
21	E Number of Bases conflict w/ Running Total	SEQ 3
21	E Wrong Nucleic Acid Designator	SEQ 3
21	E Wrong Nucleic Acid Designator	SEQ 3

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CORRECTION SUMMARY
PATENT APPLICATION US/08/631,470A

DATE: 11/17/1998
TIME: 18:03:11

Input Set: **H631470A.RAW**

Line Original Text

Corrected Data

21 SEQ 3

seq 3